\_\_\_\_\_\_

Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2009; month=4; day=25; hr=20; min=44; sec=34; ms=741; ]

\_\_\_\_\_\_

Reviewer Comments:

<210> 3

<211> 40

<212> DNA

<213> primer for S. uberis dna

The above <213> response is invalid, per Sequence Rules. The only valid responses are: the Genus species of the organism, "Artificial Sequence," or "Unknown." "Artificial Sequence" and "Unknown" require explanation in the <220>-<223> section; please give the source of the genetic material. Same error in Sequence 4.

\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*

## Validated By CRFValidator v 1.0.3

Application No: 10524198 Version No: 4.0

Input Set:

Output Set:

**Started:** 2009-04-16 09:54:53.972

**Finished:** 2009-04-16 09:54:54.339

**Elapsed:** 0 hr(s) 0 min(s) 0 sec(s) 367 ms

Total Warnings: 4

Total Errors: 0

No. of SeqIDs Defined: 6

Actual SeqID Count: 6

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W	213	Artificial or Unknown found in <213> in SEQ ID (5)
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<110> Nuijten, Petrus J. Hensen, Selma M. <120> Streptococcus Uberis Protein, Nucleic Acid Sequence Encoding the same and its use a Mastitis Vaccine <130> 2002.013 US <140> 10524198 <141> 2005-02-10 <150> EP 02078325.4 <151> 2002-08-12 <150> PCT/EP2003/008704 <151> 2003-08-06 <160> 6 <170> PatentIn version 3.3 <210> 1 <211> 603 <212> DNA <213> Streptococcus uberis <220> <223> Chromosomal DNA <400> 1 atgtttaaat tittaaagcg tgttgttitt ctagcttttc tgattitttg titttatcaa 60 gcttatataa cacatcaaaa tgtacaaaat gtcatgcaat acaaaccaat ggttgaaaaa 120 accttggctg aaaatgatac gactgccaat gtcaatttag ttttagcaat gatctacaca 180 gaaacaaaag gtggtcaggc agatgtcatg caatctagcg aaagtagtag tggtgtgact 240 aactcaatta ccgacagtca atctagtatt caacacggtg tcaaactctt gtctgagaat 300 ttgactttag ctgagaaagc tggagtagac tcttggactg cagtacaagc ttacaatttt 360 ggaacagctt acattgatta tgtggcaaaa aatggtggtg acaacactat ctctttggct 420 agtcattatt ctaaaagtgt tgtagctcca agtttaggga ataaggatgg aaaaatgtat 480 540 ttatattacc atccaattgc cctcctctat ggcggtaaac tttatcaaaa tggtggtaat atttattatt cacgagaagt tcattttaat tattacctca tacaattatt atctaaattt 600

603

taa

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Gln Tyr Lys Pro Met Val Glu Lys Thr Leu Ala Glu Asn Asp Thr Thr
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His His His His 50